SCORE Search Results Details for Application 10540215 and Search Result 20070413_115700_us-10-540-215-3.rge.

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This page gives you Search Results detail for the Application 10540215 and Search Result 20070413_115700_us-10-540-215-3.rge.

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OM nucleic - nucleic search, using sw model

Run on: April 14, 2007, 02:47:24 ; Search time 1159 Seconds

(without alignments)

1371.561 Million cell updates/sec

Title: US-10-540-215-3

Perfect score: 23

Sequence: 1 cactgaggggagaggactggggt 23

Scoring table: IDENTITY NUC

Gapop 10.0 , Gapext 1.0

Searched: 7568541 seqs, 34560148153 residues

Total number of hits satisfying chosen parameters: 15137082

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 300 summaries

Database : GenEmbl:*

1: qb env:*

2: qb pat:*

3: gb ph:*

4: gb pl:*

5: gb_pr:*

6: ·gb_ro:*

7: gb_sts:*

8: gb_sy:*

9: gb_un:*

10: gb_vi:*

11: gb_ov:*

12: gb_htg:*

13: gb_in:*

14: gb_om:*

15: gb_ba:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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4	23	100.0	861	2	CQ840731	CQ840731 Sequence
5	23	100.0	861	2	CQ840732	CQ840732 Sequence
6	23	100.0	861	2	CQ971472	CQ971472 Sequence
7	23	100.0	861	2 .	CQ971473	CQ971473 Sequence
8	23	100.0	861	2	CQ971474	CQ971474 Sequence
9	23	100.0	893	2	AR134541	AR134541 Sequence
10	23	100.0	893	2	AR134542	AR134542 Sequence
11	23	100.0	893	2	AR279531	AR279531 Sequence
12	23	100.0	893	2	AR279532	AR279532 Sequence
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               (bases 1 to 698)
REFERENCE
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            Policastro, P.F., Daniels-McQueen, S., Carle, G. and Boime, I.
  AUTHORS
  TITLE
            A map of the hCG beta-LH beta gene cluster
  JOURNAL
            J. Biol. Chem. 261 (13), 5907-5916 (1986)
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 AUTHORS
            Bellet, D., Bidart, J.-M., Vidaud, M. and Lazar, V.
  TITLE
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REFERENCE
 AUTHORS
            Zimmermann, G. and Alexander, H.
 TITLE
            Method and means for determining specific conditions or changes in
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 TITLE
           Method and means for determining specific conditions or changes in
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REFERENCE
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Bellet, D., Bidart, J.-M., Vidaud, M. and Lazar, V.

AUTHORS

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            Strausberg, R.L., Feingold, E.A., Grouse, L.H., Derge, J.G.,
  AUTHORS
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            Butterfield, Y.S., Krzywinski, M.I., Skalska, U., Smailus, D.E.,
            Schnerch, A., Schein, J.E., Jones, S.J. and Marra, M.A.
            Generation and initial analysis of more than 15,000 full-length
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            Proc. Natl. Acad. Sci. U.S.A. 99 (26), 16899-16903 (2002)
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  AUTHORS
            Strausberg, R.
  TITLE
            Direct Submission
  JOURNAL
            Submitted (13-DEC-2002) National Institutes of Health, Mammalian
            Gene Collection (MGC), Cancer Genomics Office, National Cancer
            Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
            USA
  REMARK
            NIH-MGC Project URL: http://mgc.nci.nih.gov
            Contact: MGC help desk
COMMENT
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: ATCC
            cDNA Library Preparation: Rubin Laboratory
            cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
            DNA Sequencing by: National Institutes of Health Intramural
            Sequencing Center (NISC),
            Gaithersburg, Maryland;
            Web site: http://www.nisc.nih.gov/
            Contact: nisc mgc@nhgri.nih.gov
            Akhter, N., Ayele, K., Beckstrom-Sternberg, S.M., Benjamin, B.,
            Blakesley, R.W., Bouffard, G.G., Breen, K., Brinkley, C., Brooks, S.,
            Dietrich, N.L., Granite, S., Guan, X., Gupta, J., Haghighi, P.,
            Hansen, N., Ho, S.-L., Karlins, E., Kwong, P., Laric, P., Legaspi, R.,
            Maduro, Q.L., Masiello, C., Maskeri, B., Mastrian, S.D., McCloskey, J.C.,
            McDowell, J., Pearson, R., Stantripop, S., Thomas, P.J., Touchman, J.W.,
            Tsurgeon, C., Vogt, J.L., Walker, M.A., Wetherby, K.D., Wiggins, L.,
            Young, A., Zhang, L.-H. and Green, E.D.
            Clone distribution: MGC clone distribution information can be found
            through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
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http://es/ScoreAccessWeb/GetItem.action?AppId=10540215&seqId=1362752&ItemName=20070413_11... 4/17/2007

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                                                       Indels
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                                                                      Gaps -
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RESULT 14
HUMCG7B2
LOCUS
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DEFINITION
           Human chorionic gonadotropin beta-subunit (CG-beta-7) gene, exon 1.
ACCESSION
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VERSION
            M13503.1 GI:180432
KEYWORDS
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SEGMENT
            2 of 2
SOURCE
            Homo sapiens (human)
 ORGANISM Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Euarchontoglires; Primates; Haplorrhini;
            Catarrhini; Hominidae; Homo.
REFERENCE
              (bases 1 to 984)
            Policastro, P.F., Daniels-McQueen, S., Carle, G. and Boime, I.
 AUTHORS
 TITLE
            A map of the hCG beta-LH beta gene cluster
  JOURNAL
            J. Biol. Chem. 261 (13), 5907-5916 (1986)
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PUBMED
            2422163
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ORIGIN
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                          100.0%; Pred. No. 49;
 Matches 23; Conservative
                                 0; Mismatches
                                                  0; Indels
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Qу
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PKDHPLTCDDPRFQASSSSKAPPPSLPSPSRLPGPSDTPILPQ" 1087. .1320

intron

/note="intron II"

1654 polyA site

/note="polyadenylation site"

ORIGIN

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